

Improving Early Risk Stratification in Sepsis Through Optimized
Blood Culture Time-to-Positivity Modeling

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THE PROBLEM

• Sepsis: ≥ 1.7 million US adults/year; $\geq 350,000$ deaths annually; the single most expensive condition in US hospital care.¹

• Bloodstream infection (BSI) is the most lethal sepsis phenotype: $\sim 575,000$ US cases/year; case fatality 15–25% overall, higher in septic shock.²

• Every hour of delay to appropriate antimicrobial therapy in septic shock raises mortality by $\sim 7.6\%$.³

THE BIOMARKER & THE GAP

• Blood culture time-to-positivity (TTP) reflects organism doubling time and bacterial burden at time of blood collection — proposed as an early prognostic biomarker.⁴

• Shorter TTP correlates with higher bacterial burden and, in pathogen-specific cohorts (e.g., *Staphylococcus aureus*), worse outcomes.⁴

• But two high-quality multicenter studies (FABLED 2021, Hamilton 2022) find TTP is NOT associated with mortality — the US/Western literature is inconsistent.^{5,6}

Confounding by empiric antibiotics: high-suspicion patients receive antibiotics before culture results (mortality mitigated); low-suspicion BSIs missed empirically suffer the full delay penalty ($\sim 7.6\%/h$, Kumar). This bidirectional effect attenuates any unadjusted TTP–mortality signal.

TWO INTERNATIONAL SOLUTIONS

• Non-linear modeling (Ljungquist 2025, Sweden):⁷ population-based cohort used restricted cubic splines to characterize the TTP–severity relationship without assuming linearity.

• mTTP correction (Tsal 2025, Taiwan):⁸ adjusted TTP for transport time in 1,015 community-onset BSI patients; mTTP outperformed raw TTP for 30-day mortality prediction.

RESEARCH QUESTIONS

- Among adult patients with a first organism-positive blood culture episode at UC Davis Health (2019–2026), does blood culture incubation time-to-positivity (TTP), modeled with 5-knot restricted cubic splines and adjusted for age and sex, demonstrate a statistically significant non-linear independent association with hospital length of stay (primary outcome) or all-cause mortality (secondary, exploratory outcome)?
- After correcting TTP for the measured pre-analytical specimen transport interval to produce modified TTP (mTTP = TTP + transport time), does prognostic performance for all-cause mortality improve relative to unadjusted TTP; and if mTTP correction fails, is this attributable to the high degree of pre-analytical transport heterogeneity at UCDH (median 120 minutes) compared with published reference cohorts (approximately 18 minutes; Tsal 2025)?
- What is the distribution of Collection-to-Incubator time (CTI) for positive blood cultures at UCDH, what proportion incur delays exceeding operationally relevant thresholds (≥ 4 hours, ≥ 8 hours), and how does this burden vary across patient care settings — emergency department, intensive care unit, inpatient ward, and external/satellite facilities? What is the contribution of the midnight-to-6 AM courier gap to this distribution?
- Does low blood volume per culture bottle — as a proxy for reduced bacterial inoculum — independently identify patients with higher all-cause mortality? Is the single-bottle pattern consistent with the inoculum hypothesis? (Qin 2026)

METHODS

Design and setting: Retrospective cohort study, University of California, Davis Health (UCDH), January 2019–May 2026. Pilot data were extracted from the electronic Health record via Slicer-Dicer. Confirmatory data extraction from the Electronic Antibiotic Decision Support (EADS) system is pending.

Participants: Index (first) organism-positive blood culture episode per patient ($n = 8,659$ first-per-patient episodes; 8,454 complete-case analytic cohort). Adults ≥ 18 years ($n = 7,570$) are the primary analysis; pediatric patients < 18 years ($n = 884$) analyzed separately per Qin 2026. Bottle-structure subgroups (adult cohort): paired aerobic/anaerobic $n = 7,443$; low blood volume $n = 269$; single-bottle (A or N only) $n = 93$.

Exposures: (1) TTP — time from BACTEC™ instrument load to organism-positive flag, in hours (receipt timestamp proxy). (2) mTTP = TTP + pre-analytical transport interval (Tsal 2025 primary exposure).

Outcomes: (1) All-cause mortality — patient vital status at EHR extract (May 24, 2026); variable follow-up 0–7 years (2) Hospital length of stay (log-transformed; $n = 6,675$ adults).

Statistical analysis: Non-linear TTP–outcome associations modeled via 5-knot restricted cubic splines (RCS). Likelihood ratio test (LRT; 3 d.f.) vs. linear TTP specification.

Covariates: age (3-knot RCS), sex. Contamination sensitivity: three-tier exclusion (laboratory PC flag; organism-based CoNS/Micrococcus/Corynebacterium; union).

PILOT vs CONFIRMATORY DATA

Pilot (this poster) — Slicer-Dicer extract:

- TTP from receipt-to-result timestamp (proxy for true BACTEC instrument-load).
- Mortality = vital status at extract (May 2026); follow-up 0–7 yr; not 30-day.
- Contamination: single-flag heuristic (not paired-set adjudicated).
- No severity scores, no antibiotic timing.

Confirmatory — EADS extraction (pending) will provide:

- Verified BACTEC load timestamps (true TTP, true mTTP).
- Registry-linked 30-day all-cause mortality.
- Per-bottle volumes for covariate-adjusted inoculum analysis.
- SOFA components, lactate, procalcitonin, CRP.
- Paired-set contamination adjudication.

— The pilot establishes pipeline + methods; EADS will refine every endpoint and adjustment.

Other caveats: TTP–mortality attenuated after laboratory-flagged contamination exclusion ($p=0.12$ – 0.20 across three-tier sensitivity; PC-flag excl. $p=0.20$, organism-based excl. $p=0.12$); TTP–LOS directionally robust but also attenuated ($p=0.077$ – 0.17). Unadjusted courier-gap mortality comparison is acutely-confounded (fast-transport cultures select for sicker patients on STAT orders).

TRANSPORT WORKFLOW (CTI)

UCDH median CTI is $\sim 7\times$ longer than published international reference cohorts.

Median CTI: **120 min**

IQR 86–184 min · Mean 169 min · P95 454 min

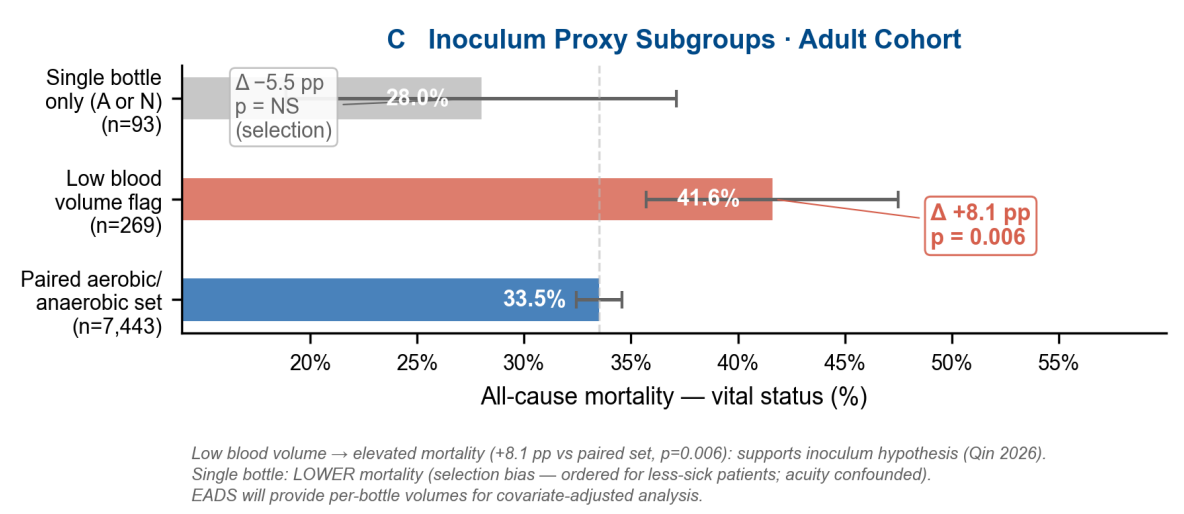
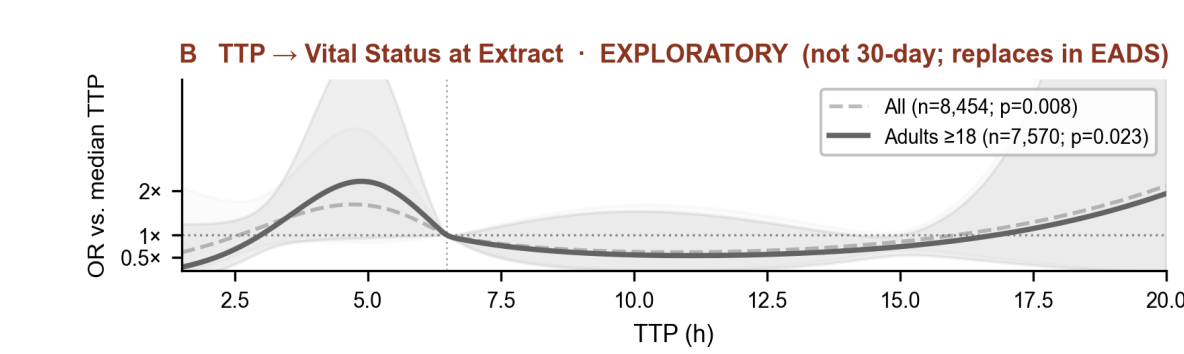
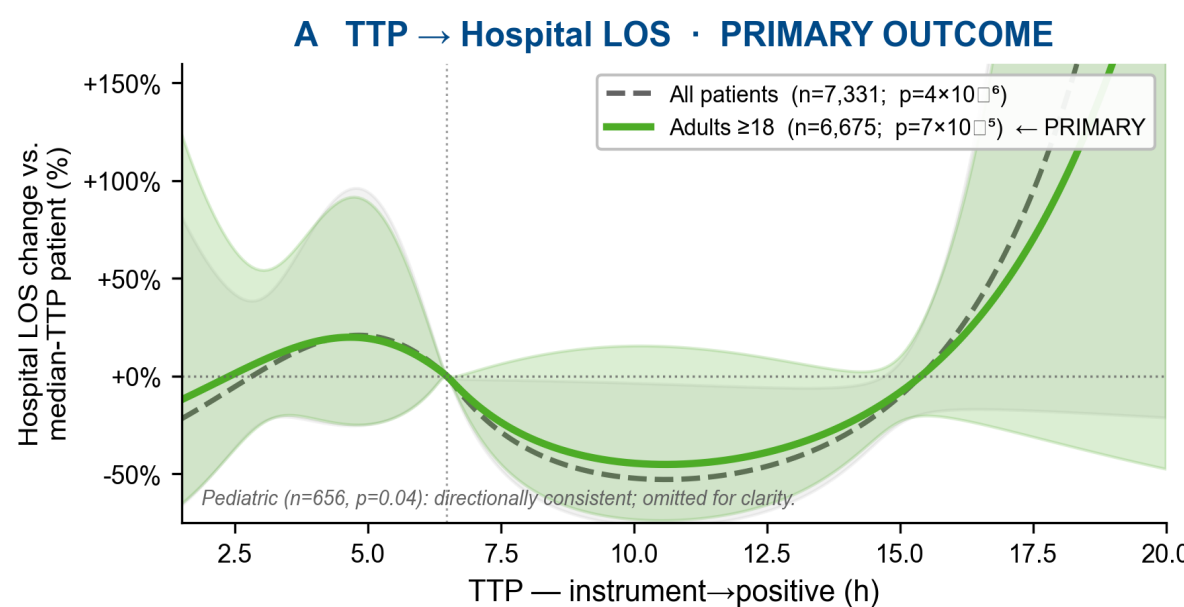
Published reference: **~ 18 min**

Tsal 2025 cohort (Taiwan)

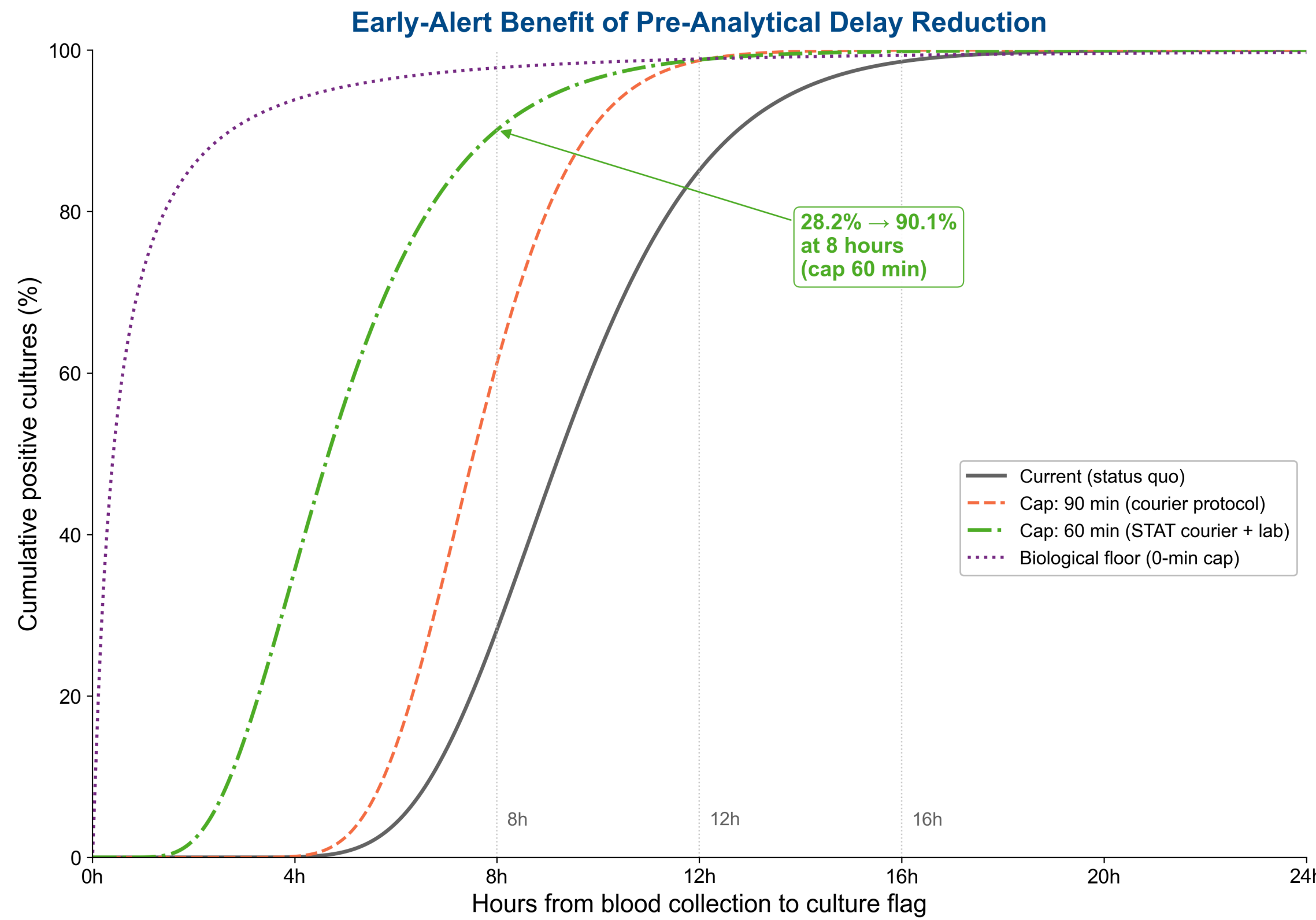
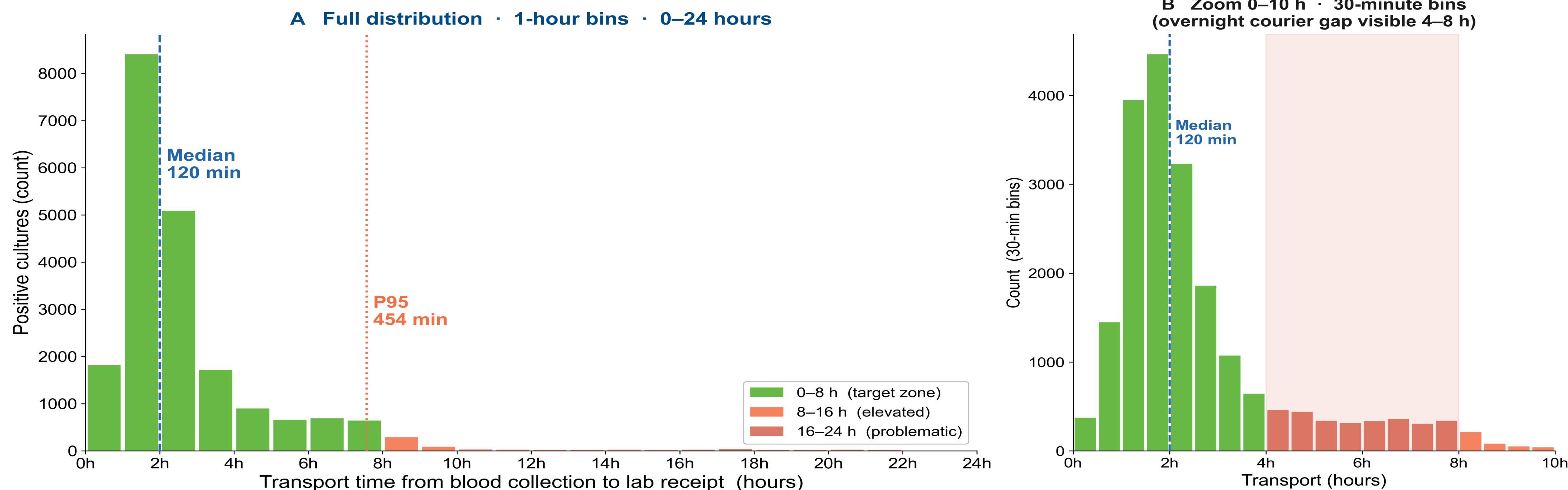
Cultures positive by 8 h: **28.2%**

logistical status quo prior to 2026 intervention

HVC-QI gap: midnight–6 AM courier blackout delays $\sim 15\%$ of cultures by 4–6 h



Pre-Analytical Transport Delay Distribution · n = 20,710 positive cultures, 2019–2026



CONCLUSIONS

» Non-linear TTP–hospital LOS is the primary robust finding (total cohort $p = 4\times 10^{-6}$; adult primary analysis $p = 7\times 10^{-6}$), driven by inpatient strata and consistent across sex and racial/ethnic groups.

» The strongest TTP–mortality association is in the intensive care unit subgroup ($p < 0.001$, $n = 976$), where earlier TTP results would be most clinically actionable for antibiotic escalation and de-escalation. The whole-cohort mortality finding is exploratory (vital-status endpoint; will be resolved in EADS).

» Modified TTP (mTTP) correction failed to improve prognostic performance at UCDH; CTI is approximately 7-fold longer than the Tsal 2025 reference setting (median 120 min vs. ~ 18 min), adding non-biological heterogeneity that attenuates the mTTP–mortality signal ($p = 0.325$).

» Low blood volume specimens had higher all-cause mortality ($+8.1$ pp, $p = 0.006$), supporting the inoculum hypothesis (Qin 2026). Single-bottle cultures had lower mortality than paired sets (-5.5 pp, NS), consistent with acuity-selection confounding rather than a biological inoculum effect — underscoring the need for per-bottle volumes in inoculum-adjusted analysis.

» The midnight-to-6 AM courier gap generates an accumulation of 4–8 h CTI delays (visible in Panel B, Figure A). Targeted overnight courier runs or STAT prioritization would directly reduce this zone and advance blood culture positivity flagging by up to 6 hours.

NEXT STEPS

• EADS extraction: verified BACTEC timestamps, registry-linked 30-day mortality, per-bottle volumes for inoculum-adjusted analysis.

• Severity covariate adjustment: SOFA score, serum lactate, procalcitonin.

• Overnight courier pilot (on-going, starting in early 2026): quantify the CTI reduction for midnight–6 AM draws at UCDH

RESULTS — ADULT COHORT

$n = 7,570$ adults · LOS subset $n = 6,675$ · Pediatric $n = 884$ (separate)

Model performance (primary analysis)

Exposure	Outcome	p (non-linear)	C / R ⁴
TTP (inst.→pos.)	log(LOS)	7.2×10^{-6} PRIMARY	R² = 0.041
TTP (inst.→pos.)	Mortality ↑	0.023 exploratory	C = 0.64
mTTP (coll.→pos.)	Mortality ↑	0.325 null	C = 0.65

† Mortality endpoint = vital status at EHR extract (May 2026); variable follow-up 0–7 yr. NOT 30-day. Will be replaced in EADS.

SUBGROUP FINDINGS

- Adult Low blood volume cohort ($n=269$): 41.8% mortality vs. paired-set reference ($n=7,443$) 33.5% ($+8.1$ percentage points; $\chi^2=7.72$, $p=0.006$). Supports the inoculum hypothesis: inadequate specimen volume may reflect harder-to-phenotomize patients or true under-sampling of the bacteremic inoculum.
- Single bottle only (A or N; $n=93$): 28.0% mortality vs. 33.5% paired (-5.5 pp; $p = NS$). Counter to the inoculum hypothesis. Most likely explanation: single-bottle cultures are ordered preferentially for less-critically ill patients (acuity selection bias), not driven by reduced inoculum. EADS per-bottle volumes will enable adjusted analysis.
- Intensive care unit (ICU subgroup ($n=976$)): 41.0% all-cause mortality vs. 30.3% non-ICU. TTP–mortality association strongest in ICU (LRT $p<0.001$, $\chi^2=23.9$).
- Pediatric cohort ($n=884$): TTP–hospital LOS directionally consistent with adults (LRT $p=0.04$). Bottle structure subgrouping is clinically distinct: recommended blood volume is weight-based (minimum 1–3 mL/kg); making standard adult paired-set criteria non-applicable. EADS will provide weight-adjusted per-bottle volumes for pediatric inoculum analysis.

BY CARE SETTING

Adult cohort mortality and TTP outcome p-values per stratum

Setting	n	Mort%	Med LOS	TTP → Mort p	TTP → LOS p
Emergency (ED)	4,138	29.7%	6 d	0.47	0.07
ICU	976	41.0%	25 d	<0.001 ★	0.05
Other inpatient	3,545	31.4%	17 d	0.18	<0.001 ★

★ ICU: TTP→mortality is the strongest subgroup signal. Other inpatient: TTP→LOS is the strongest subgroup signal.

EQUITY — OUTCOME MODEL (TTP → Hospital LOS, adult cohort)

5-knot restricted cubic spline TTP to linear TTP specification, adjusted for age and sex. All strata: longer TTP associated with longer hospital LOS.

Stratum	n (LOS subset)	LRT $\chi^2(3)$	p value	Interpretation
Male	4,186	21.8	<0.001	Non-linear TTP→LOS confirmed; consistent with primary adult model
Female	2,677	11.1	0.011	Non-linear TTP→LOS confirmed; consistent with primary adult model
White	3,408	13.6	0.004	Non-linear TTP→LOS confirmed; consistent with primary adult model
Other / Unknown	1,864	14.7	0.002	Non-linear TTP→LOS confirmed; consistent with primary adult model
Asian	637	7.7	0.053	Borderline significance ($p = 0.053$); statistical power limited
Black / African American	954	5.1	0.17	Directionally consistent; not significant; limited stratum power ($n = 954$)

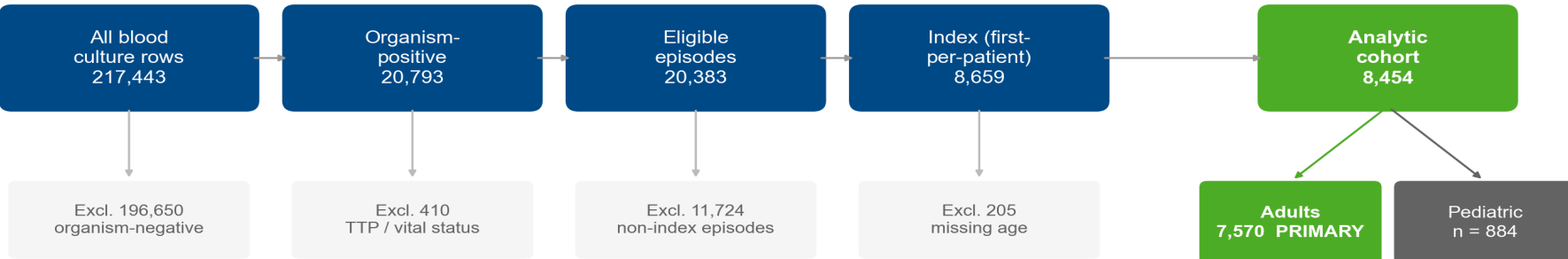
EQUITY — PRE-ANALYTICAL TRANSPORT BY SETTING

Collection → lab receipt, all cultures 2019–2026 ($n = 207,630$ with valid timestamp)

Setting	N	Median	IQR	≥ 4 h (%)
ED	86,143	110 min	80–162 min	14%
ICU (adult)	26,834	112 min	83–166 min	15%
Inpatient ward	69,605	127 min	92–184 min	16%
PICU / NICU	8,555	118 min	87–176 min	15%
External/satellite	10,672	315 min	201–533 min	67%

Within-hospital settings comparable (110–127 min). External/satellite (Vibra, dialysis, regional labs): 315 min median, 3× longer, 67% ≥ 4 h. These patients are older and have higher comorbidities. Satellite courier optimization is the highest-leverage workflow target.

COHORT FLOW



DEMOGRAPHICS (Table 1)

Variable	Adults (n = 7,570)	Pediatric (n = 884)
Age, yr. median [IQR]	62 [48–72]	1 [0–9]
Male sex	4,612 (60.9%)	525 (59.4%)
Race: White	3,765 (49.6%)	—
Race: Black / African American	1,045 (13.8%)	—
Race: Asian	659 (8.7%)	—
Race: Other / Unknown	2,111 (27.9%)	—
Mortality †	2,497 (33.0%)	125 (14.1%)
Hospital LOS ‡ med [IQR]	16 [8–23] d	—

† Vital status at EHR extract (May 2026); variable follow-up 0–7 yr; ‡ LOS available for 6,675 of 7,570 adults (88.2%).

mTTP — SUPPLEMENTARY

mTTP (collection→positive) null for mortality (adult $p=0.325$). Heterogeneous transport drowns the mTTP signal.

Measure	LRT p	Direction	Interpretation
TTP (incubation)	0.023	Non-linear	Shorter TTP → longer LOS / higher mortality; primary robust finding
mTTP (coll.→pos.)	0.325	Null	UCDH CTI adds variance (120 min med. vs. ref. ~ 18 min; Tsal 2025)

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